FIG. 1A

1	CCTAGAGCCAGCAGAGTCCAGGCTGCTGTTAACAACTTCATGTCCCCGTGGGTAGCAGGC	60
61	${\tt AGGTGCTTCTGATCTGGCTCTCCTTGACCACTGTACTCATCAAATAGACCAAGATC}$	120
121	CCCAGAGTCCAAGATCCTTACAAGGGGGCCAGAAAGGGATGAGCTTTCTGAAGAAGCACT	180
181	GATGTAAAATACCAGGAATTTTGACATCGAAGAAGATTTTTGTGATGGCAGCTGGGATTT	240
241	GGCCATAATCTAGAAGACACATGGTGAATACAGTTGCAAGTCATTTAGTCATATTTCTTG	300
301	CTAAATTGCTGTGTCTTCAATGGCTGAATTGAAGATCCCTCTTACCCGCCAGGTGCCAAG	360
361	AACTATGAACAGGCAGGCAATAGAAAGACAACTAAAGAAGGATCCAACGATTTGAAATT	420
421	CCAGAACTTCAGTCTGCCAAAAAACAGGTCATGGCCTCGCATCAATAGTGCCACAGGCCA	480
481	GTACCAGAGGATGAACAAGCCTCTTCTAGACTGGGAAAGAAA	540
541	TGGAGCAAAAGGCCACAGTGATGATGACTATGATGACCCTGAGCTTCGGATGGAAGAGAC	600
601	ATGGCAGTCGATTAAAATTTTACCAGCCCGGCCTATAAAGGAATCTGAATATGCAGATAC	660
661	${\tt ACACTATTTCAAGGTTGCAATGGACACTCCCCTTCCGTTAGACACCAGGACCTCTATCTC}$	720
721	CATTGGACAGCCGACCTGGAACACACAGACGAGGTTGGAAAGAGTGGACAAACCCATTTC	780
781	CAAGGACGTCAGAAGCCAAAACATTAAAGGAGATGCATCCGTAAGAAAGA	840
841	TTTACCACCTCCTCGGCCTCTCATAACACTTCCGAAGAAGTACCAACCCTTGCCCCCTGA	900
901	GCCGGAGAGCAGCCACCTTTATCTCAGAGACACCCTTTCCAGAAGTCCAGAGAAT	960
961	GCCCAGTCAGATAAGCTTAAGGGACTTAAGTGAGGTCCTTGAAGCAGAAAAAGTTCCTCA	1020
1021	TAACCAGAGGAAGCCTGAATCAACTCATCTGTTAGAAAACCAAAATACTCAAGAGATTCC	1080
1081	ACTTGCCATTAGCAGTTCTTCATTCACGACAAGCAACCACAGTGTGCAAAACAGAGATCA	1140
1141	TAGAGGAGGCATGCAGCCCTGTTCTCCTCAGAGATGCCAGCCTCCAGCCAG	1200
1201	TCACGAAAATATACTGCCCTATAAATACACAAGCTGGAGACCACCTTTCCCCAAAAGGTC	1260
1261	TGATAGAAAGGATGTCCAGCACAATGAATGGTACATTGGAGAATACAGCCGCCAGGCAGT	1320
1321	GGAAGAGGCATTCATGAAGGAGAACAAGGATGGTAGTTCTTGGTCCGAGATTGTTCCAC	1380
1381	AAAATCCAAGGAAGAGCCCTATGTTTTGGCTGTTTTTATGAGAACAAAGTCTACAATGT	1440
1441	${\tt AAAAATCCGCTTCCTGGAGAGGAATCAGCAGTTTGCCCTGGGGACAGGACTCAGAGGAGA}$	1500

FIG. 1B

1501	$\tt TGAGAAGTTTGATTCAGTAGAAGACATCATCGAACACTACAAGAATTTTCCCATTATACT$	1560
1561	AATTGATGGGAAAGATAAAACTGGGGTCCACAGGAAACAGTGTCACCTCACTCA	1620
1621	$\tt CCCTCTCACCAGACACCTCTTGCCTCTGTAGCCTGGTCTTTGTGTTATCTTTGGTTTACT$	1680
1681	$\tt GGATTCAGCGCTTCCATTGTTTTCATTGATTTCAAAAGTTTATTTTCTGTGCCTTCAAGG$	1740
1741	GACAACTTTTTTAACTTTGGAGAAAAGAAAACACTCTATAACAGAGAGTGGAAAATCAC	1800
1801	ΤΓΑΓΕ ΕΤΤΤΤΕΔΑ ΑΕΤΤΓΑ ΑΑΓΓΑ ΓΑ ΓΑ ΓΑ ΓΑ ΑΑΑΑ ΑΑΤΑΤΤΤΑ ΤΑ ΑΓΑΤΕΓΑ ΑΑΑ	1851

FIG. 2

1							M	A	E	L	K	I	P	L	Т	R	Q	V	Ρ	R	14
15	Т	M	N	R	Q	G	N	R	K	Т	Т	K	Ε	G	S	N	D	L	K	F	34
35	Q	N	F	s	L	P	K	N	R	S	W	P	R	I	N	S	A	Т	G	Q	54
55	Y	Q	R	M	N	K	P	L	L	D	W	E	R	N	F	Α	A	V	L	D	74
75	G	A	K	G	Н	S	D	D	D	Y	D	D	P	E	L	R	M	Ε	Ε	Т	94
95	W	Q	S	I	K	I	L	P	A	R	P	I	K	E	S	E	Y	A	D	Т	114
115	Н	Y	F	K	V	A	M	D	T	P	L	P	L	D	T	R	Т	s	I	S	134
135	I	G	Q	P	Т	M	N	Т	Q	Т	R	L	E	R	V	D	K	P	I	S	154
155	K	D	V	R	S	Q	N	I	K	G	D	A	S	V	R	K	N	K	I	P	174
175	L	Р	P	Р	R	P	L	I	Т	L	P	K	K	Y	Q	P	L	Р	P	E	194
195	P	E	S	S	R	P	P	L	S	Q	R	Н	Т	F	P	E	V	Q	R	M	214
215	P	S	Q	I	S	L	R	D	L	S	E	V	L	E	A	E	K	V	P	Н	234
235	N	Q	R	K	P	E	S	Т	Н	L	L	Ε	N	Q	N	Т	Q	Ε	I	P	254
255	L	А	I	S	S	S	S	F	Т	Т	S	N	Н	S	V	Q	N	R	D	Н	274
275	R	G	G	M	Q	P	С	S	Р	Q	R	С	Q	P	P	A	S	С	S	P	294
295	Н	E	N	I	L	P	Y	K	Y	Т	S	M	R	Р	Ρ	F	F	K	R	. S	314
315	D	R	K	D	V	Q	Н	N	Ε	M	Y	I	G	Е	Y	S	R.	Q	A	V	334
335	E	E	А	F	М	K	E	N	K	D	G	S	F	L	V	R	D	С	5	T	354
355	K	S	K	Ε	Ε	P	Y	V	L	A	V	F	Y	E	N	K	V	Y	. 1	ı v	374
375	K	I	R	F	L	E	R	N	Q	Q	F	A	L	G	T	' G	· I	, R		; D	394
395	E	K	F	D	S	V	Ε	D	I	I	Ε	Н	Y	K	N	T F	P	r			414
415	I	D	G	K	D	K	Т	G	V	Н	R	K	: Ç) C	. H	I L	ı 7.	. Č) I	? L	
435	P	· L	T	R	Н	L	L	P	L	ı											443

FIG. 3A

1	CCTAGAGCCAGCAGAGTCCAGGCTGCTGTTAACAACTTCATGTCCCCGTGGGTAGCAGGC	60
61	AGGTGCTTCTGTCTGATCTGGCTCTCCTTGACCACTGTACTCATCAAATAGACCAAGATC	120
121	CCCAGAGTCCAAGATCCTTACAAGGGGGCCAGAAAGGGATGAGCTTTCTGAAGAAGCACT	180
181	GATGTAAAATACCAGGAATTTTGACATCGAAGAAGATTTTTGTGATGGCAGCTGGGATTT	240
241	GGCCATAATCTAGAAGACACATGGTGAATACAGTTGCAAGTCATTTAGTCATATTTCTTG	300
301	CTAAATTGCTGTGTCTTCAATGGCTGAATTGAAGATCCCTCTTACCCGCCAGGTGCCAAG M A E L K I P L T R Q V P R	360 14
361 15	AACTATGAACAGGCAGGCAATAGAAAGACGACTAAAGAAGGATCCAACGATTTGAAATT T M N R Q G N R K T T K E G S N D L K F	420 34
421 35	CCAGAACTTCAGTCTGCCAAAAAACAGGTCATGGCCTCGCATCAATAGTGCCACAGGCCA Q N F S L P K N R S W P R I N S A T G Q	480 54
481 55	GTACCAGAGGATGAACAAGCCTCTTCTAGACTGGGAAAGAAA	540 74
541 75	TGGAGCAAAAGGCCACAGTGATGATGACTATGATGACCCTGAGCTTCGGATGGAAGAGAC G A K G H S D D D P E L R M E E T	600 94
601 95	T ATGGCAGTCGATTAAAATTTTACCAGCCCGGCCTATAAAGGAATCTGAATATGCAGATAC W Q S I K I L P A R P I K E S E Y A D T	660 114
661 115	ACACTATTTCAAGGTTGCAATGGACACTCCCCTTCCGTTAGACACCAGGACCTCTATCTC H Y F K V A M D T P L P L D T R T S I S	720 134
	CATTGGACAGCCGACCTGGAACACACAGACGAGGTTGGAAAGAGTGGACAAACCCATTTC	780
721 135	I G Q P T W N T Q T R L E R V D K P I S	154
781 155	CAAGGACGTCAGAAGCCAAAACATTAAAGGAGATGCATCCGTAAGAAAGA	840 174
841	TTTACCACCTCCTCGGCCTCTCATAACACTTCCGAAGAAGTACCAACCCTTGCCCCCTGA	900
175	<i>LPPPRP</i> LITLPKKYQ <i>PLPPE</i>	194
901 195	GCCGGAGAGCAGCACCTTTATCTCAGAGACACACCTTTCCAGAAGTCCAGAGAAT $m{p}$ E S S R P P L S Q R H T F P E V Q R M	960 214
961 215	GCCCAGTCAGATAAGCTTAAGGGACTTAAGTGAGGTCCTTGAAGCAGAAAAAGTTCCTCA PSQISLRDLSEVLEAEKVPH	1020 234
1021 235	TAACCAGAGGAAGCCTGAATCAACTCATCTGTTAGAAAAACCAAAATACTCAAGAGATTCC NQRKPESTHLLENQNTQEIP	1080 254
1081 255	ACTTGCCATTAGCAGTTCTTCATTCACGACAAGCAACCACAGTGTGCAAAACAGAGATCA L A I S S S F T T S N H S V O N R D H	114(274
1141	TAGAGGAGGCATGCAGCCTGTTCTCCTCAGAGATGCCAGCCTCCAGCCAG	1200

FIG. 3B

275	R	G	G	М	Q	P	C	S	P	Q	R	С	Q	P	P	Α	S	C	S	P	294
1201	TCA	.CGA	AAA	TAT	'ACT	GCC	CTA	TAA	ATA	CAC	AAC	CTG	GAG	ACC	ACC	TTT	'CCC	CAA	AAG	GTC	1260
295	H	E	N	I	L	P	Y	K	Y	T	S	W	R	₽	P	F	₽	К	R	S	314
1261	TGA	TAC	AAA	.GGA	TGI	CCA	GCA	CAA	TGA	ATO	GTA	CAT	TGG	AGA	ATA	CAG	CCG	CCA	.GGC	AGT	1320
315	D	R	K	D	V	Q	H	N	E	W	Y	I	G	Ε	Y	s	R	Q	A		334
1321	GGA	AGA	\GGC	'ATI	CAT	'GAA	AGGA	GAA	CAA	AGGA	TGG	OATE	TTT	CTT	GGT	CCG	AGA	TTG	TTC	CAC	1380
335	_E	E	A	F	M	K	E	N	K	D	G	S	F	L	V	R	D	C	S	T	354
1381	AAA	ATC	CAA	GGA	AGA	GCC	CTA	TGI	TTT	GGC	TGI	GTI	TTP	TGA	GAZ	ACAA	AGT	'C'TA	CAA	TGT	1440
355	_K	S	K	E	E	P	Y		L	A	V	F	Y	E	N	K	V	<u>Y</u>	N	V	374
1441	AAA	LAA	CCC	CTI	CCT	'GGA	GAG	GAZ	ATCA	AGCA	GTI	TGC	CCI	'GGG	GAC	CAGG	ACI	'CAG	AGG	AGA	1500
375	K	I	R	F	L	E	R	N	Q	Q	F	A	L	G	T	G	L	R	G	D	394
1501	TGA	GAA	GTI	TGA	TTC	AGT	AGA	AGA	ACAT	CAT	CGA	AACA	CTA	CAA	GAA	TTT	TCC	CAT	'TAT	ACT	1560
395	E	K	F	D	S	V	E	D	I	I	Ε	H	Y	K	N	F	P	I	I	L	414
1561	AAT	'TGA	ATGO	GAA	AGA	AATA	AAA	TGG	GGT	CCF	CAC	GAA	ACA	GTG	TCA	ACCI	CAC	TCA	.GCC	'ACT	1620
415	I	D	G	K	D	K	T	G	V	H	R	K	Q	С	H	L	\mathbf{T}	Q	P	L	434
1621	CCC	TCT:	CAC	CAC	ACA	CCI	CTI	GCC	CTCI	GTA	\GC(TGG	TCT	TTC	TGT	TAT	CTT	'T'GG	TTT	ACT	1680
435	P	L	T	R	H	L	L	P	L	*											444
1681	GGA	TTC	CAGC	GCT:	TCC	ATI	GTT	TTC	CATT	GAT	TTC	CAAZ	AAGI	TTF	TTT	TCT	GTG	CCI	TCA	AGG	1740
1741	GAC	'AAC	TTT	TTT.	'AAC	CTTT	GG <i>I</i>	AGAZ	AAA	AAF	AAA	CACT	CTF	ATAF	ACAC	GAGA	\GTG	GAA	ΓAΑ	CAC	1800
1801	TCA	CGG	TTT	TGA	AAC	TTC	CAA	ACCE	CAC	SAGI	AAA	TAT	TT	TAZ	ACA1	rgca	AAA	L			1851

FIG. 4A

60	GTCAGACCTCTCAGGTCTGTGGCTGCATTTCACAGGAAACCAAGTCTAAAAACGGACCTAT	1
120	CAGGAGGTTTTCTGCTGAAGGGCACTGCTTAGCATCGAGAAGAATTCAACCCACCGCCTT	61
180	ACTAATTTCCAGTGCCCCAAGGTCTCTGCACTGCCGCCCCTCCTCACAGGAGACGGACAC	121
240	$\tt CTCAGCCTAGATCCCTTGGTGCTCTCCACGCTGTTCAGGCTGAATTGAAGATCCCTCTTA$	181
300	CCCGCCAGGTGCCAAGAACTATGAACAGGCAGGGCAATAGAAAGACAACTAAAGAAGGAT	241
360	CCAACGATTTGAAATTCCAGAACTTCAGTCTGCCAAAAAACAGGTCATGGCCTCGCATCA	301
420	ATAGTGCCACAGGCCAGTACCAGAGGATGAACAAGCCTCTTCTAGACTGGGAAAGAAA	361
480	TTGCTGCAGTCCTGGATGGAGCAAAAGGCCACAGTGATGATGACTATGATGACCCTGAGC	421
540	TTCGGATGGAAGAGACATGGCAGTCGATTAAAATTTTACCAGCCCGGCCTATAAAGGAAT	481
600	CTGAATATGCAGATACACACTATTTCAAGGTTGCAATGGACACTCCCCTTCCGTTAGACA	541
660	CCAGGACCTCTATCTCCATTGGACAGCCGACCTGGAACACACAGACGAGGTTGGAAAGAG	601
720	TGGACAAACCCATTTCCAAGGACGTCAGAAGCCAAAACATTAAAGGAGATGCATCCGTAA	661
780	GAAAGAACAAGATTCCTTTACCACCTCCTCGGCCTCTCATAACACTTCCGAAGAAGTACC	721
840	AACCCTTGCCCCCTGAGCCGGAGAGCAGCAGCCACCTTTATCTCAGAGACACACCTTTC	781
900	CAGAAGTCCAGAGAATGCCCAGTCAGATAAGCTTAAGGGACTTAAGTGAGGTCCTTGAAG	841
960	CAGAAAAAGTTCCTCATAACCAGAGGAAGCCTGAATCAACTCATCTGTTAGAAAACCAAA	901
1020	ATACTCAAGAGATTCCACTTGCCATTAGCAGTTCTTCATTCA	961
1080	TGCAAAACAGAGATCATAGAGGAGGCATGCAGCCCTGTTCTCCTCAGAGATGCCAGCCTC	1021
1140	CAGCCAGCTGCAGCCCTCACGAAAATATACTGCCCTATAAATACACAAGCTGGAGACCAC	1081
1200	CTTTCCCCAAAAGGTCTGATAGAAAGGATGTCCAGCACAATGAATG	1141
1260	ACAGCCGCCAGGCAGTGGAAGAGGCATTCATGAAGGAGAACAAGGATGGTAGTTTCTTGG	1201
1320	TCCGAGATTGTTCCACAAAATCCAAGGAAGAGCCCTATGTTTTGGCTGTGTTTTATGAGA	1261
1380	ACAAAGTCTACAATGTAAAAATCCGCTTCCTGGAGAGGAATCAGCAGTTTGCCCTGGGGA	1321
1440	CAGGACTCAGAGGAGATGAGAAGTTTGATTCAGTAGAAGACATCATCGAACACTACAAGA	1381

FIG. 4B

1441	ATTTTCCCATTATACTAATTGATGGGAAAGATAAAACTGGGGTCCACAGGAAACAGTGTC	1500
1501	ACCTCACTCAGCCACTCCCTCTCACCAGACACCTCTTGCCTCTGTAGCCTGGTCTTTGTG	1560
1561	${\tt TTATCTTTGGTTTACTGGATTCAGCGCTTCCATTGTTTTCATTGATTTCAAAAGTTTATT}$	1620
1621	${\tt TTCTGTGCCTTCAAGGGACAACTTTTTTAACTTTGGAGAAAAAGAAAAACACTCTATAACA}$	1680
1681	GAGAGTGGAAAATCACTCACGGTTTTGAAAGTTCAAACCACAGAGAAAATATTTATAACA	1740
1741	TGCAAAAATAAAAACATTCTAGTAACTGGCCACTGGAAAATAAAT	1800
1801	GGGTTTTAAAAGTATCTTCTAAAAAACAACAACAAAAAATACTATAAACATAGCCATTAT	1860
1861	GCTCATGATACAGGCGAGCAGCAAAGGGCACCAGAAGCTGTTGCTTAAATGTTTGCAGTC	1920
1921	AGTGCAAGACAAGTCTATGGGAAATTCCCAAATCTGTGCTCTTTACAGGACACTGCGCTG	1980
1981	${\tt CCTTTATGTCAGTTGTTGGGCCTTACATATATACAATGTGTGGATGATTTCTTACACTAA}$	2040
2041	AGATGCTGGGCTGCGGTGCCTCATGCCTGTAATCCCAGCACTTTGGGAGGCTGAGG	2100
2101	TGGACAGATCACGAGGTCAGGAGATCAAGACCATCCTGGCTAACATGGTGAAACCCCCATG	2160
2161	TCTACTAAAAATACAAAAATCAGCTGGGCGTGGTGGTGGGTG	2220
2221	TCGGGAGGCTGAGGCAGGAGAATGGTGTGAACCCCGGGAGGCGGAGCTTGCAGTGAGCCGA	2280
2281	AATCGCGCCACTGCACTCCAATCCAGCCTGGGGACAGAGAGACTCCGTCTCAAAA	2335

FIG. 5

1							M	N	R	Q	G	N	R	K	T	Т	K	Ε	G	S	14
15	N	D	L	K	F	Q	N	F	S	L	P	K	N	R	S	W	P	R	I	N	34
35	S	A	Т	G	Q	Y	Q	R	M	N	K	P	L	L	D	W	Ε	R	N	F	54
55	A	A	V	L	D	G	A	K	G	Н	S	D	D	D	Y	D	D	P	Ε	L	74
75	R	M	E	E	Т	M	Q	S	I	K	I	L	P	A	R	Р	I	K	E	S	94
95	E	Y	A	D	Т	Н	Y	F	K	V	A	M	D	Т	P	L	P	L	D	Т	114
115	R	Т	S	I	S	I	G	Q	P	Т	M	N	Т	Q	Т	R	L	E	R	V	134
135	D	K	P	I	S	K	D	V	R	S	Q	N	I	K	G	D	A	S	V	R	154
155	K	N	K	I	P	L	P	P	P	R	Р	L	I	Т	L	P	K	K	Y	Q	174
175	P	L	P	P	E	P	E	S	S	R	Р	P	L	S	Q	R	Н	Т	F	P	194
195	E	V	Q	R	M	P	S	Q	I	S	L	R	D	L	S	Ε	V	L	Ε	A	214
215	Ε	K	V	P	Н	N	Q	R	K	P	Ε	S	Т	Н	L	L	E	N	Q	N	234
235	Т	Q	E	I	P	L	Α	I	S	S	S	S	F	T	Т	S	N	Н	S	V	254
255	Q	N	R	D	Н	R	G	G	M	Q	P	С	S	P	Q	R	С	Q	P	Ρ	274
275	А	S	С	S	P	Н	E	N	I	L	P	Y	K	Y	Т	S	W	R	P	P	294
295	F	P	K	R	S	D	R	K	D	V	Q	Н	N	Ε	W	Y	I	G	E	Y	314
315	S	R	Q	A	V	Ε	E	A	F	M	K	E	N	K	D	G	S	F	\mathbf{L}	V	334
335	R	D	С	S	Т	K	S	K	E	E	P	Y	V	L	A	V	F	Y	Ε	N	354
355	K	V	Y	N	V	K	I	R	F	L	E	R	N	Q	Q	F	A	L	G	Т	374
375	G	L	R	G	D	Ε	K	F	D	S	V	Ε	D	I	I	Ε	Н	Y	K	N	394
395	F	P	I	I	L	I	D	G	K	D	K	Т	G	V	Н	R	K	Q	С	Н	414
415	L	Т	Q	P	L	Р	L	Т	R	Н	L	L	Р	L							428

FIG. 6A

1	GTCAGACCTCTCAGGTCTGTGGCTGCATTTCACAGGAAACCAAGTCTAAAACCGACCTAT	60
61	CAGGAGGTTTTCTGCTGAAGGGCACTGCTTAGCATCGAGAAGAATTCAACCCACCGCCTT	120
121	ACTAATTTCCAGTGCCCCAAGGTCTCTGCACTGCCGCCCCTCCTCACAGGAGACGGACAC	180
181	CTCAGCCTAGATCCCTTGGTGCTCTCCACGCTGTTCAGGCTGAATTGAAGATCCCTCTTA	240
241 1	CCCGCCAGGTGCCAAGAACTATGAACAGGCAGGCAATAGAAAGACAACTAAAGAAGGAT M N R Q G N R K T T K E G S	300 14
301	CCAACGATTTGAAATTCCAGAACTTCAGTCTGCCAAAAAACAGGTCATGGCCTCGCATCA	360
15	N D L K F Q N F S L P K N R S W P R I N	34
361	ATAGTGCCACAGGCCAGTACCAGAGGATGAACAAGCCTCTTCTAGACTGGGAAAGAAA	420
35	SATGQYQRMNKPLLDWERNF	54
421	TTGCTGCAGTCCTGGATGGAGCCAAAAGGCCACAGTGATGATGACTATGATGACCCTGAGC	480
55	AAVLDGAKGHSDDDYDDPEL	74
481	TTCGGATGGAAGAGACATGGCAGTCGATTAAAATTTTACCAGCCCGGCCTATAAAGGAAT	540
75	RMEETWQSIKILPARPIKES	94
541	CTGAATATGCAGATACACACTATTTCAAGGTTGCAATGGACACTCCCCTTCCGTTAGACA	600
95	EYADTHYFKVAMDTPLPLDT	114
601	CCAGGACCTCTATCTCCATTGGACAGCCGACCTGGAACACACAGACGAGGTTGGAAAGAG	660
115	RTSISIGQPTWNTQTRLERV	134
661	TGGACAAACCCATTTCCAAGGACGTCAGAAGCCAAAACATTAAAGGAGATGCATCCGTAA	720
135	D K P I S K D V R S Q N I K G D A S V R	154
721	GAAAGAACAAGATTCCTTTACCACCTCCTCGGCCTCTCATAACACTTCCGAAGAAGTACC	780
155	KNKIPLPPRRPLITLPKKYQ	174
781	AACCCTTGCCCCCTGAGCCGGAGAGCAGCAGCCACCTTTATCTCAGAGACACACCTTTC	840
175	PLPPESSRPPLSQRHTFP	194
841	CAGAAGTCCAGAGAATGCCCAGTCAGATAAGCTTAAGGGACTTAAGTGAGGTCCTTGAAG	900
195	EVQRMPSQISLRDLSEVLEA	214
901	CAGAAAAAGTTCCTCATAACCAGAGGAAGCCTGAATCAACTCATCTGTTAGAAAACCAAA	960
215	EKVPHNQRKPESTHLLENQN	234
961	ATACTCAAGAGATTCCACTTGCCATTAGCAGTTCTTCATTCA	
235	TQEIPLAISSSSFTTSNHSV	254
1021	TGCAAAACAGAGATCATAGAGGAGGCATGCAGCCCTGTTCTCCTCAGAGATGCCAGCCTC	1080
255	Q N R D H R G G M Q P C S P Q R C Q P P	274
1081	CAGCCAGCTGCAGCCCTCACGAAAATATACTGCCCTATAAATACACAAGCTGGAGACCAC	1140
	ASCSPHENILPYKYTSWRPP	294

FIG. 6B

141	CTTTCCCCAAAAGGTCTGATAGAAAGGATGTCCAGCACAATGAATG	1200
295	F P K R S D R K D V Q H N E W Y I G E Y	314
L201	ACAGCCGCCAGGCAGTGGAAGAGGCATTCATGAAGGAGAACAAGGATGGTAGTTTCTTGG	1260
315	S R Q A V E E A F M K E N K D G S F L V	334
1261	TCCGAGATTGTTCCACAAAATCCAAGGAAGAGCCCTATGTTTTGGCTGTGTTTTATGAGA	1320
335	R D C S T K S K E E P Y V L A V F Y E N	354
1321	ACAAAGTCTACAATGTAAAAATCCGCTTCCTGGAGAGGAATCAGCAGTTTGCCCTGGGGA	1380
355	KVYNVKIRFLERNQQFALGT	374
	CAGGACTCAGAGGAGATGAGAAGTTTGATTCAGTAGAAGACATCATCGAACACTACAAGA	1440
1381	G L R G D E K F D S V E D I I E H Y K N	394
375	G L K G D F K L D 2 4 F D 1 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	
1441	ATTTTCCCATTATACTAATTGATGGGAAAGATAAAACTGGGGTCCACAGGAAACAGTGTC	1500
395	FPIILIDGKDKTGVHRKQCH	414
1501	ACCTCACTCAGCCACTCCCTCTCACCAGACACCTCTTGCCTCTGTAGCCTGGTCTTTGTG	1560
415	LTQPLPLTRHLLPL	429
1561	TTATCTTTGGTTTACTGGATTCAGCGCTTCCATTGTTTTCATTGATTTCAAAAGTTTATT	1620
1601	TTCTGTGCCTTCAAGGGACAACTTTTTTAACTTTGGAGAAAAGAAAACACTCTATAACA	1680
1621	11C1G1GCC11CAAGGGACAC111111AAG111GGAGAALLAGALLAG	
1681	GAGAGTGGAAAATCACTCACGGTTTTGAAAGTTCAAACCACAGAGAAAATATTTATAACA	1740
1741	TGCAAAAATAAAACATTCTAGTAACTGGCCACTGGAAAATAAAT	1800
1801	GGGTTTTAAAAGTATCTTCTAAAAAACAACAACAAAAAATACTATAAACATAGCCATTAT	1860
1861	GCTCATGATACAGGCGAGCAGCAAAGGGCACCAGAAGCTGTTGCTTAAATGTTTGCAGTC	1920
1921	AGTGCAAGACAAGTCTATGGGAAATTCCCAAATCTGTGCTCTTTACAGGACACTGCGCTG	1980
1981	${\tt CCTTTATGTCAGTTGTTGGGCCTTACATATATACAATGTGTGGATGATTTCTTACACTAA}$	2040
2041	AGATGCTGGGCTGGGTGCCTCATGCCTGTAATCCCAGCACTTTGGGAGGCTGAGG	2100
2101	TGGACAGATCACGAGGTCAGGAGATCAAGACCATCCTGGCTAACATGGTGAAACCCCCATG	2160
2161	TCTACTAAAAATACAAAAAATCAGCTGGGCGTGGTGGTGGCTGCCTGTAGTCCCAGCTAC	2220
2221	TCGGGAGGCTGAGGCAGGAGAATGGTGTGAACCCGGGAGGCGGAGCTTGCAGTGAGCCGA	2280
2281	AATCGCGCCACTGCACTCCAATCCAGCCTGGGGACAGAGAGACTCCGTCTCAAAA	2335

FIG. 7A

1	GGCTGCTGTTAACAACTTCATGTCCCCGTGGGTAGCAGGCAG	80
61	GCTCTCCTTGACCACTGTACTCATCAAATAGACCAAGATCCCCAGAGTCCAAGATCCTTA	120
121	CAAGGGGCCCAGAAAGGGATGAGCTTTCTGAAGAAGCACTGATGTAAAATACCAGGAATT	180
181	TTGACATCGAAGAAGATTTTTGTGATGGCAGCTGGGATTTGGCCATAATCTAGAAGACAC	240
241	ATGGTGAATACAGTTGCAAGTCATTTAGTCATATTTCTTGCTAAATTGCTGTGTCTTCAA	300
301	TGGGGCAATAGAAAGACTAAAGAAGGATCCAACGATTTGAAATTCCAGAACTTCAGT	360
361	CTGCCAAAAAACAGGTCATGGCCTCGCATCAATAGTGCCACAGGCCAGTACCAGAGGATG	420
421	AACAAGCCTCTTCTAGACTGGATTTGGCAGCTTGACCATTTATTATCGCACAGTGGATGC	480
481	AATCAGAAGTCTGGGCACAGCATGGCTCAACTAGTTCCCCTGTTCTGGGTCTCACAAGAC	540
541	TGAAAGCAACATGCTGGCAGGGCTGCATTCTCCTCCAGGGGCTCTGAAGAGGAACTTGCT	600
601	TCCAGATTCTTTCAGGAAAGAAACTTTGCTGCAGTCCTGGATGGA	660
661	GATGATGACTATGATGACCCTGAGCTTCGGATGGAAGAGACATGGCAGTCGATTAAAATT	720
721	TTACCAGCCCGGCCTATAAAGGAATCTGAATATGCAGATACACACTATTTCAAGGTTGCA	780
781	ATGGACACTCCCCTTCCGTTAGACACCAGGACCTCTATCTCCATTGGACAGCCGACCTGG	840
841	AACACACAGACGAGGTTGGAAAGAGTGGACAAACCCATTTCCAAGGACGTCAGAAGCCAA	900
901	AACATTAAAGGAGATGCATCCGTAAGAAAGAACAAGATTCCTTTACCACCTCCTCGGCCT	960
961	CTCATAACACTTCCGAAGAAGTACCAACCCTTGCCCCCTGAGCCGGAGAGCAGCAGCCA	1020
1021	CCTTTATCTCAGAGACACACCTTTCCAGAAGTCCAGAGAATGCCCAGTCAGATAAGCTTA	1080
1081	AGGGACTTAAGTGAGGTCCTTGAAGCAGAAAAAGTTCCTCATAACCAGAGGAAGCCTGAA	1140
1141	TCAACTCATCTGTTAGAAAACCAAAATACTCAAGAGATTCCACTTGCCATTAGCAGTTCT	1200
1201	TCATTCACGACAAGCAACCACAGTGTGCAAAACAGAGATCATAGAGGAGGCATGCAGCCC	1260
1261	TGTTCTCCTCAGAGATGCCAGCCTCCAGCCAGCTGCAGCCCTCACGAAAATATACTGCCC	1320
1323	1 TATAAATACACAAGCTGGAGACCACCTTTCCCCAAAAGGTCTGATAGAAAGGATGTCCAG	1380
120	CACAATGAATGGTACATTGGAGAATACAGCCGCCAGGCAGTGGAAGAGGCATTCATGAAG	144

FIG. 7B

1441	GAGAACAAGGATGGTAGTTTCTTGGTCCGAGATTGTTCCACAAAATCCAAGGAAGAGCCC	1500
1501	${\tt TATGTTTTGGCTGTGTTTTATGAGAACAAAGTCTACAATGTAAAAATCCGCTTCCTGGAG}$	1560
1561	${\tt AGGAATCAGCAGTTTGCCCTGGGGACAGGACTCAGAGGAGATGAGAAGTTTGATTCAGTA}$	1620
1621	GAAGACATCATCGAACACTACAAGAATTTTCCCATTATACTAATTGATGGGAAAGATAAA	1680
1681	ACTGGGGTCCACAGGAAACAGTGTCACCTCACTCAGCCACTCCCTCTCACCAGACACCTC	1740
1741	$\tt TTGCCTCTGTAGCCTGGTCTTTGTGTTATCTTTGGTTTACTGGATTCAGCGCTTCCATTG$	1800
1801	${\tt TTTTCATTGATTTCAAAAGTTTATTTTCTGTGCCTTCAAGGGACAACTTTTTTAACTTTG}$	1860
1861	GAGAAAAGAAAACACTCTATAACAGAGAGTGGAAAATCACTCAC	1920
1921	AACCACAGAGAAAATATTTATAACATGCAAAAAATAAAAACATTCTAGTAACTGGCCACT	1980
1981	GGAAAATAAAAAAAAAACTAGGGTTTTAAAAGTATCTTCTAAAAAACAACAACAA	2040
2041	AAAATACTATAAACATAGCCATTATGCTCATGATACAGGCGAGCAGCAAAGGGCACCAGA	2100
2101	AGCTGTTGCTTAAATGTTTGCAGTCAGTGCAAGACAAGTCTATGGGAAATTCCCAAATCT	2160
2161	GTGCTCTTTACAGGACACTGCGCTGCCTTTATGTCAGTTGTTGGGCCCTTACATATATACA	2220
2221	ATGTGTGGATGATTTCTTACACTAAAGATGCTGGGCTGG	2280
2281	ATCCCAGCACTTTGGGAGGCTGAGGTGGACAGATCACGAGGTCAGGAGATCAAGACCATC	2340
2341	CTGGCTAACATGGTGAAACCCCCATGTCTACTAAAAATACAAAAAATCAGCTGGGCGTGGT	2400
2401	GGTGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGTGTGAACCCG	2460
2461	GGAGGCGGAGCTTGCAGTGAGCCGAAATCGCGCCACTGCACTCCAATCCAGCCTGGGGAC	2520
2521	AGAGAGACTCCGTCTCAAAA	2540

FIG. 8

1											M	Ε	Ε	Т	M	Q	S	I	K	I	10
11	L	P	Α	R	P	I	K	E	S	E	Y	A	D	Т	Н	Y	F	K	V	A	30
31	M	D	Т	P	L	P	L	D	Т	R	Т	S	I	S	I	G	Q	P	Т	W	50
51	N	Т	Q	Т	R	L	Ε	R	V	D	K	P	I	S	K	D	V	R	S	Q	70
71	N	I	K	G	D	А	S	V	R	K	N	K	I	P	L	P	₽	P	R	P	90
91	L	I	Т	L	P	K	K	Y	Q	P	L	P	P	Ε	P	E	S	S	R	P	110
111	P	L	S	Q	R	Н	Т	F	P	E	V	Q	R	M	P	S	Q	I	S	L	130
131	R	D	L	S	Е	V	L	E	А	Ε	K	V	P	Н	N	Q	R	K	P	E	150
151	s	Т	Н	L	L	Ε	N	Q	N	Т	Q	Е	I	Р	L	A	I	S	S	S	170
171	S	F	Т	Т	S	N	Н	S	V	Q	N	R	D	Н	R	G	G	M	Q	P	190
191	С	s	P	Q	R	С	Q	P	P	A	S	С	S	Ρ	Н	E	И	I	L	P	210
211	Y	K	Y	Т	S	W	R	P	P	F	Ρ	K	R	S	D	R	K	D	V	Q	230
231	Н	N	Ε	W	Y	I	G	E	Y	S	R	Q	A	V	Ε	E	A	F	M	K	250
251	E	N	K	D	G	S	F	Ŀ	V	R	D	С	S	Т	K	S	K	Ε	E	Р	270
271	Y	V	L	A	V	F	Y	E	N	K	V	Y	N	V	K	I	R	F	L	E	290
291	R	N	Q	Q	F	A	L	G	Т	G	L	R	G	D	E	K	F	D	S	V	310
311	E	D	I	I	E	Н	Y	K	N	F	Р	I	I	L	I	D	G	K	D	K	330
331	Т	G	V	Н	R	K	Q	С	Н	L	Т	Q	P	L	P	L	T	R	Н	L	350
351	L	Р	L																		353

FIG. 9A

1	GGCTGCTGTTAACAACTTCATGTCCCCGTGGGTAGCAGGCAG	60													
61	GCTCTCCTTGACCACTGTACTCATCAAATAGACCAAGATCCCCAGAGTCCAAGATCCTTA														
121	CAAGGGGGCCAGAAAGGGATGAGCTTTCTGAAGAAGCACTGATGTAAAATACCAGGAATT														
181	TTGACATCGAAGAAGATTTTTGTGATGGCAGCTGGGATTTGGCCATAATCTAGAAGACAC	240													
241	ATGGTGAATACAGTTGCAAGTCATTTAGTCATATTTCTTGCTAAATTGCTGTCTCTCAA	300													
301	TGGGGCAATAGAAAGACTAAAGAAGGATCCAACGATTTGAAATTCCAGAACTTCAGT	360													
361	CTGCCAAAAAACAGGTCATGGCCTCGCATCAATAGTGCCACAGGCCAGTACCAGAGGATG	420													
421	AACAAGCCTCTTCTAGACTGGATTTGGCAGCTTGACCATTTATTATCGCACAGTGGATGC														
481	AATCAGAAGTCTGGGCACAGCATGGCTCAACTAGTTCCCCTGTTCTGGGTCTCACAAGAC														
541	$\tt TGAAAGCAACATGCTGGCAGGGCTGCATTCTCCTCCAGGGGCTCTGAAGAGGAACTTGCT$														
601	TCCAGATTCTTTCAGGAAAGAAACTTTGCTGCAGTCCTGGATGGA	660													
661 1	GATGATGACTATGATGACCCTGAGCTTCGGATGGAAGAGACATGGCAGTCGATTAAAATT MEET WQSIKI	720 10													
721 11	TTACCAGCCCGGCCTATAAAGGAATCTGAATATGCAGATACACACTATTTCAAGGTTGCA L P A R P I K E S E Y A D T H Y F K V A	780 30													
781 31	ATGGACACTCCCCTTCCGTTAGACACCAGGACCTCTATCTCCATTGGACAGCCGACCTGG M D T P L P L D T R T S I S I G Q P T W	840 50													
841 51	AACACACAGACGAGGTTGGAAAGGGACGACAAACCCATTTCCAAGGACGTCAGAAGCCAANT QTRLERVDKPISKDVRSQ	900 70													
901 71	AACATTAAAGGAGATGCATCCGTAAGAAAGAACAAGATTCCTTTACCACCTCCTCGGCCT N I K G D A S V R K N K I P L P P P R P	960 90													
961 91	CTCATAACACTTCCGAAGAAGTACCAACCCTTGCCCCCTGAGCCGGAGAGCAGCAGCCA L I T L P K K Y Q P L P P E P E S S R P	1020 110													
1021 111	CCTTTATCTCAGAGACACCCTTTCCAGAAGTCCAGAGAATGCCCAGTCAGATAAGCTTA PLSQRHTFPEVQRMPSQISL	1080 130													
1081 131		1140 150													
1141 151	TCAACTCATCTGTTAGAAAACCAAAATACTCAAGAGATTCCACTTGCCATTAGCAGTTCT S T H L L E N Q N T Q E I P L A I S S S	1200 170													
1201 171	TCATTCACGACAAGCAACCACAGTGTGCAAAACAGAGATCATAGAGGAGGCATGCAGCCC S F T T S N H S V Q N R D H R G G M Q P	1260 190													
1261	TGTTCTCCTCAGAGATGCCAGCCTCCAGCCAGCTGCAGCCCTCACGAAAATATACTGCCC	1320													

FIG. 9B

191	С	S	P	Q	R	С	Q	P	P	A	S	С	S	P	H	E	N	I	L	P	210
1321										TTT F	CCC P	CAA K	AAG R	GTC S	TGA D	TAG. R	AAA K	GGA D	TGT V	CCAG O	1380 230
211	Y	K	Y	Т	S	W	R	Ρ	P	_	_		-						·	~	250
1381	CA																			GAAG	1440
231	Η	N	E	W	Y	I	G	Ε	Y	S	R	Q	A	V	Ε	E	A	F	M	K	250
1441	GA	.GAA	.CAA	.GGA	TGG	TAG	TTT	CTT	GGT	CCG	AGA	TTG	TTC	CAC	!AAA	ATC	CAA	GGA	AGA	GCCC	1500
251	Ε	N	K	D	G	S	F	L	V	R	D	С	S	Т	K	S	K	Ε	Ε	P	270
1501	TA	TGT	TTT	'GGC	'TGT	GTT	'TTA	TGA	.GAA	CAA	AGT	'CTA	.CAA	TGT	'AAA	AAT	CCG	CTT	CCT	GGAG	1560
271	Y	V	L	A	V	F	Y	E	N	K	V	Y	N	V	K	I	R	F	L	E	290
1561	70.00	ור א א	man.	aa 7	CTT	TOO	יכפת	ccc	ca c	יז ככ	лст	יראמ	יז ממ	מאמא	тсъ	CDD	GTT	тса	ጥጥር	AGTA	1620
1561 291	R	AADI N	O	0 0	F	A	L L	G	T	.AGG	AC1 L	R	G	D	E	K	F	D	S	V	310
			~	~																	
1621																				TAAA	1680 330
311	Ε	D	Ι	Ι	Ε	H	Y	K	N	F	Ρ	Ι	I	L	I	D	G	K	D	K	330
1681	AC	TGG	GGT	CCA	CAG	GAA	ACA	GTG	TCA	CCT	'CAC	TCA	.GCC	ACT	CCC	TCT	'CAC	CAG	ACA	.CCTC	1740
331	\mathbf{T}	G	V	H	R	K	Q	C	H	L	\mathbf{T}	Q	P	L	P	L	\mathbf{T}	R	H	L	350
4544				ama		aa	man		man	1017 70 00	ama	maa	ımmı	17 CIT	ימטזי	THE C	יזארייר	ימפיי	יידיכו	ATTG	1800
1741 351	Tu L	.GCC P	TCT L	.GT.Z *	AGCC	TGC	TCI	TT.G	ilGl	IAI	CII	. LGC	1 T T T	AC I	.GGF	TIL	.AGC	.60 !	100	AIIG	354
331																					
1801	TT	TTC	TTA	'GA'I	TTC	'AAA	AGT	TTF	TTT	TCT	GTO	CCI	TCF	AAGG	GAC	AAC	TTT:	TTT	'AAC	TTTG	1860
1861	CI 7	א מים	7770	ኋ አ አ ፖ	\ 7\ 7\ C	י א רייז	ערוים	ממידי	ער אַ כּ	:DCD	CTC.		רעע	ייים	יייכו	יכפפ	יייי	ጥርል	AAG	TTCA	1920
1001	GF	MA	MA	J.C.Y.C.Y.C	m	.AC I	CIF	1171	10110	Jrior.	.0			. 0110	,						
1921	A	ACCA	ACAG	BAGA	AAA	TAT	TTA	TAF	CAT	GCA	AAA	raa/	'AAZ	AAA	CATI	CTA	GTA	ACT	'GGC	CACT	1980
1981	CC	ית א <i>ד</i> אדי	ילידי ער	יי אל אל א	רי א א ד	ייאאי	אר ארי	. א א ר	יייי א כ	יממיז	ուհուհուհ	מ מ מי	۱ ک <i>ا</i> کا ۱	ים מי	المنابات	ነጥ አ ጀ	<u> </u>	ACA	מטמג	ACAA	2040
1901	GC	3 <i>1</i> -71-71-	747 F	7.7.7.	LAAF	75-75-7	. AAA	MAC	LAC	1001	1	LWW	mo.		,,,,					10.11	
2041	AZ	LAA	ACT	TAT	AAA	CATA	AGCC	CTA	ATO	CTC	ATC	BATA	CAC	GCC	BAG	CAGC	AAA	\GG@	CAC	CAGA!	2100
0101	70.0	n Cimc	ımm <i>c</i>	acmo	ר אום	man		1070	יייייי	(CITIC	1 C T T	\ C 7\ C	ግ አ አረ	ī m 🖰 a	ቦ አጥር	יממז	דיממי	ייייייי	ממטי	ATCT	2160
2101	AC	JC 10	31.10	-C11	LAAA	11G1	LIIC	CAC	3 I CF	7010	CAF	MUMC	AA	J1C.	LAIC	JUUL	זבייני ז	. 1 0 1	·Cru	MICI	2100
2161	G:	rgc1	rct1	CTAC	CAGO	BAC	ACTO	GCGC	CTGC	CTT	TAT	rgtc	CAG	rtg:	rtgo	GCC	TTF	ACA'I	ATA	TACA	2220
				~ ~ ~~	~ ~ ~	-ma		1 7 CI	C 3 70 7			naaa	· ama	700	na aa	тата	יממת	יי ער איי	7000	ייייטייי א	2280
2221	A'.	rgro	3TGC	3A'1'(3A'1"I	L"I'C".	L.T.VC	JAC".	ĽAAŁ	AGA'I	.GC	rGGC	3C.T.C	تىنى	rGCC	ידטנ	i(()	.CA1	. GCC	TGTA	2200
2281	A.	rcco	CAG	CAC	rtt	GG <i>I</i>	AGGC	CTGA	AGG:	rggi	ACAC	3ATC	CAC	GAG	3TC	AGGI	AGA7	CA	AGAC	CCATC	2340
															ל מכ אד א		77.00	ma	1000	тпаат	2400
2341	C'.	rggc	CTAA	ACA'	ľGG'.	ľĠAÆ	AACC	JCCA	7.T.G.	rc'r <i>f</i>	AC.T.	AAAA	₹₩.T.¥	ACAA	AAAA	AA.T.C) DA	TGG	الالال	TGGT	2400
2401	G	GTG(GTC	3CC	rgt <i>i</i>	AGT	CCC	AGC"	rac'	rcgo	3GA(GGC1	rga(GGC2	AGG	\GAZ	ATGO	TG'	rga <i>i</i>	ACCCG	2460
0.1	~	~~~	700	~ ~ ·	~m~	7/77	יחמי		707	, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,	700	7007	v (Jun.	7/77/	ama,	71 77 77 17	יממז	\ C.C.	יידירי	rcan c	2520
2461	G()ÐAt)ئان ن)ÐAt	C.T.,T.(JCA(J L G₽	4GC(_GA!	-747.T.(الالالا	JCC#	-1C 17	JCAU		_F1F1	LCCF	3.J.C.	- 1 GC	EGGAC	اکرے
2521	A	GAG	AGA	CTC	CGT	CTC	AAAA	Ŧ.													2540

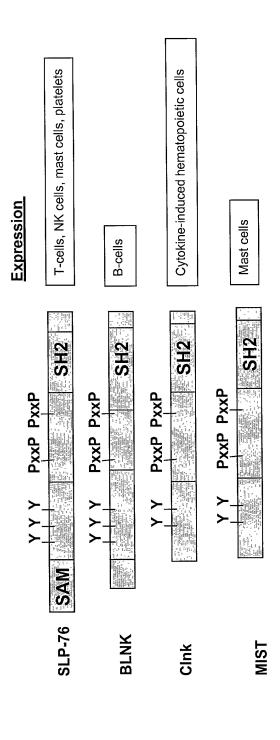


FIG. 10

FIG. 11
Recombinant MIST proteins

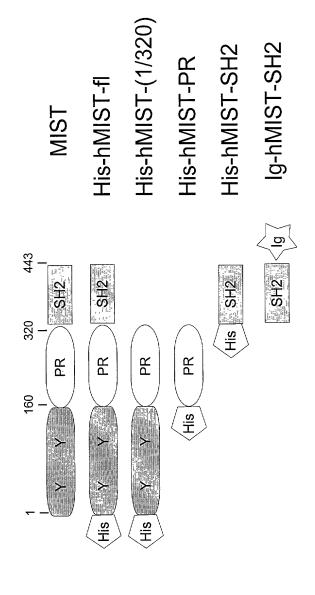


FIG. 12

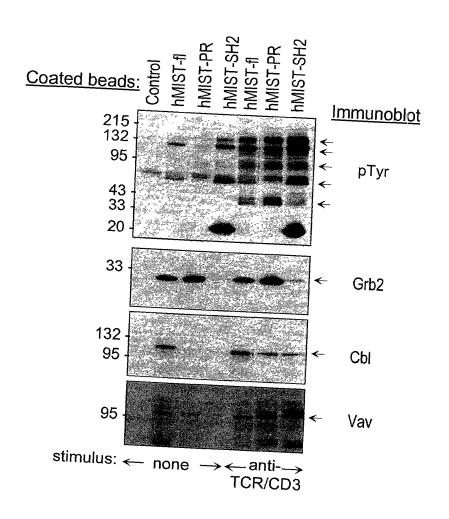
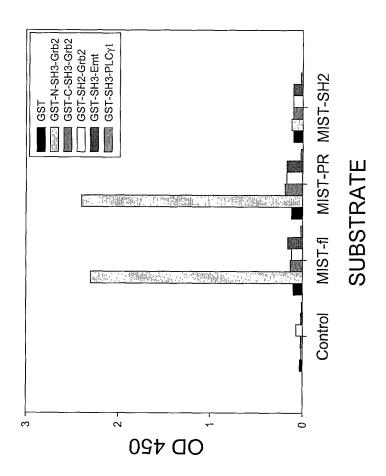


FIG. 13
Interaction between hMIST and Grb2-SH3 domain



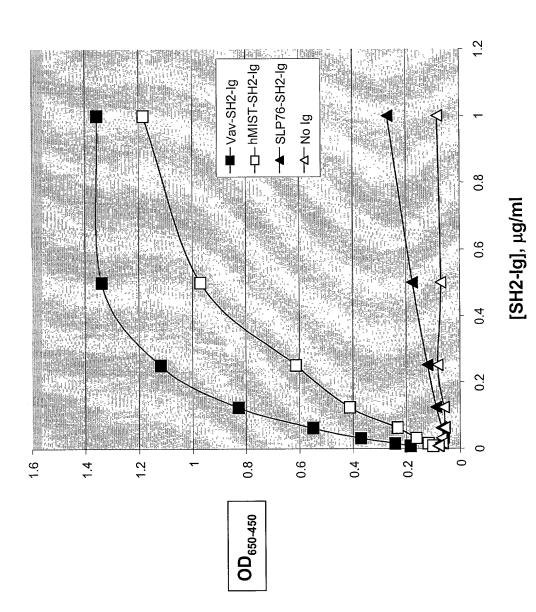


FIG. 14

